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Reviewer: Durreshwar Anjum

Timestamp: Wed Oct 10 13:45:14 EDT 2007

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Reviewer Comments:

<210> SEQ ID NO 1
<211> LENGTH: 1512
<212> TYPE: DNA
<213> ORGANISM: Ipomoea batatas
<400> SEQUENCE: 1

Please do not insert alpha-numeric headings. Also check for the
subsequent errors throughout the sequences.

Application No: 10597945 Version No: 1.0

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Output Set:

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Finished: 2007-09-24 22:12:18.027
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Total Errors: 0
No. of SeqIDs Defined: 30
Actual SeqID Count: 30

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W 213	Artificial or Unknown found in <213> in SEQ ID (29)
W 213	Artificial or Unknown found in <213> in SEQ ID (30)

<110> APPLICANT: Korea Research Institute of Bioscience and Biotechnology
KWAK, Sang-Soo
KWON, Suk-Yoon
LEE, Haeng-Soon
RYU, Sun-Hwa
<120> TITLE OF INVENTION: Multiple stress-inducible peroxidase promoter derived from Ipomoea batatas
<130> FILE REFERENCE: 58049-00035
<140> CURRENT APPLICATION NUMBER: US/10/597,945A
<141> CURRENT FILING DATE: 2006-08-11
<150> PRIOR APPLICATION NUMBER: PCT/KR2004/000381
<151> PRIOR FILING DATE: 2004-02-24
<160> NUMBER OF SEQ ID NOS: 30
<170> SOFTWARE: KopatentIn 1.71

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gtattatctg cagggttgc acgcaggctc tctactaaac gatacaccta ctttcacccgg	480
agaacagacc gccggcggca ataataactc agtcagaggt tttgaggtga tacaacaagc	540
taaagagaat gtgataacca aatgtcccta catacaagta tcttgcgcg acatcttac	600
cattgctgcc cgtgattttt tccagagatc aagtccattt atttctaaag gttgaaat	660
ataagaacaa gaatccaaac aaataacaga cagtaaaaaaa aaaagattta tttgggtttt	720
caatatgttgc aaattttttt tatatttaat gactgttattt tatgcattat atttatatgc	780
aactctaaac atgcagttt ctggagaaac gtacaccgtg actctggaa gactcgatgc	840
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aattagaaca aattaaatct cagtttgctt tattatatta ttatcaacaa taataattta	180
atactgatcg aagaacttcc ctttcaagt tctcttattt aggaaggctg agaagccatt	240
aatcctcatc atcagctcga ccactcattt cttcttcata cttccttgc tgtgataatc	300
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aaatagtact ccaatcataa taaattatct tatattatat tgccaacaat taaaaattca	180
aattagaaca aattaaatct cagtttgctt tattatatta ttatcaacaa taataattta	240
atactgatcg aagaacttcc ctttcaagt tctcttattt aggaaggctg agaagccatt	300
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atcatggtaa ggtgtttat cgtggcagca tgagtgcattg acaaacgcatttatttattt 180
taaaacaaaaa tagtactcca atcataataa attatcttat attatattgc caacaattaa 240
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taatttaataa ctgatcgaag aactttccct ttcaagttct ctatattttaagg aagcctgaga 360
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tgcttttagca gcaaaaataa gaatggtaact atacttaatc ctcatacatag tcttcaaccc 240
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tctatatttg atgatggtaa ttatgtatcat ggtaagggtgt tttatcgtgg cagcatgagt 540
gcatgacaaa cgcataatattt attataaaaa caaaatagta ctccatcat aataaattat 600
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catttaatta aatttctatc ctcatgttga acccattaat cgaataattg acatattaga 180
taaacttagc catcatatga catttgcattca tgattgtatca tttttaaaaaa ataaaaaacaa 240
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ctgtacaata ataattttgt ttcaactaag aggatgttgg caaaagtata attaaacttg 360
tgatcttcgt acaataatttca tgcttcacgc actcaactag tcacatctt ccaggcaaaa 420
tttacttttc tatgaatatg agaagttcca tctatggaaa taacggattt tttatctaat 480
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ttaatttagt catgataata caactcacct actccattat agccgataat acaactcacc 720
tagctactcc attatagttcc aacaatataca aatgaataaaa atagtaatgg tgacttaaag 780
ggctgaatcc aacatataattt ctgacattt aaaaatgttca cgtacggtaa gatttagtata 840
atgaaataaaa gttaatcatt ctctatattt gatgtatggta attagtatca tggtaaggtg 900

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acaaaattaaa	tctcagttt	cttattata	ttattatcaa	caataataat	ttaatactga	1080
tgcagaact	ttcccttca	agttctctat	ttaaggaagc	ctgagaagcc	attaatcctc	1140
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 <222> LOCATION: (1)..(1467)
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ataaaaaata	ttgggtcaaa	ataccttacc	gattttccc	aaatattcac	ggaacttact	180
gccagaatct	accctgcctt	ttcccttcac	tattttcaca	actataagca	tatatggca	240
taaatatgac	atgaacatgc	atgaaccaat	gcagggtgaa	agtaagattg	aatatactga	300
tactacaatt	aactaatgat	aaagtataac	ttttgtaaaa	aatttgattt	tttttttiga	360
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aattatgtaa	accctgtaat	ctagtaatct	gtacaataat	aattttgttt	caactaagag	600
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ataatttaat	actgatcgaa	gaactttccc	tttcaagttc	tctatTTAG	gaagcctgag	1860
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22
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